This Page Is Inserted by IFW Operations and is not a part of the Official Record

BEST AVAILABLE IMAGES

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images may include (but are not limited to):

- BLACK BORDERS
- TEXT CUT OFF AT TOP, BOTTOM OR SIDES
- FADED TEXT
- ILLEGIBLE TEXT
- SKEWED/SLANTED IMAGES
- COLORED PHOTOS
- BLACK OR VERY BLACK AND WHITE DARK PHOTOS
- GRAY SCALE DOCUMENTS

IMAGES ARE BEST AVAILABLE COPY.

As rescanning documents will not correct images, please do not report the images to the Image Problem Mailbox.



Atty. Docket No. CRP-008DV

IN THE UNITED STATES PATENT AND TRADEMARK

APPLICANT(S): Huston, James S. et al.

KAMINIER OF WILM A

OFFICE

CORRES. AND MAI

SERIAL NO.:

07/661,070

GROUP NO.: 185

FILED:

February 26, 1991

TITLE: PRODUCT AND PROCESS FOR THE PRODUCTION, ISOLATION

AND PURIFICATION OF RECOMBINANT POLYPEPTIDES

which is a DIVISIONAL OF: 07/462,297, FILED: 12/28/89 which is a CONTINUATION OF:07/028,484, FILED: 3/20/87

Honorable Commissioner of Patents & Trademarks
Washington, D.C. 20231

DECLARATION OF DAVID C. RUEGER, MARC F. CHARETTE,

CHARLES M. COHEN, ROBERTO CREA, JAMES S. HUSTON,

PETER C. KECK, HERMANN OPPERMANN, AND RICHARD J. RIDGE

UNDER 35 U.S.C. RULE 1.131

We declare:

1. We invented the subject matter claimed in the abovereferenced application, which claims priority as a divisional
application from USSN 07/462,297, filed 12/28/89, which in turn
is a continuation of 07/028,484 filed 3/20/87. We, Drs. Rueger,
Charette, Cohen, Huston, Keck, and Oppermann, are employees of
the named assignee, Creative BioMolecules, Inc. I, Dr. Ridge, am
a former employee of Creative BioMolecules, Inc. I, Dr. Crea, am
a former employee and a current member of the Board of Directors
of Creative BioMolecules, Inc.

- 2 -

- 2. Prior to March 28, 1986, we reduced to practice in the United States the construction and expression of recombinant DNA encoding a fusion polypeptide containing a leader/hinge region, a cleavage site, and a target polypeptide, and the subsequent cleavage of that polypeptide at the cleavage site by a cleavage site-specific protease. The following experiments, performed by Zita Babickas under the direction of David C. Rueger, demonstrate that a fusion polypeptide having the structure described in the above-captioned patent application is cleaved at the cleavage site nearest the hinge region. These experiments are described in the above-captioned patent application (see Example III, in particular) and in the accompanying Exhibits A and B. The results are shown in Exhibits C-E. These experiments involve fusion polypeptides in which the target protein is calcitonin.
- 3. Polypeptide 1, described on pages 39-41 of the above-captioned patent application, includes a TRP-LE leader sequence connected at its carboxyl end to a Phe-Pro-Gly hinge, followed by a glutamic acid residue and human calcitonin. As stated in the specification on page 40, lines 1-7, it was desired that the tripeptide hinge would promote cleavage at the Glu residue by providing a flexible residue (glycine) and uncharged "spacer" residues next to the Glu residue. The large leader was used to promote high expression levels. The sequence of polypeptide 1 is provided in the specification on pages 40-41.

- 4. Polypeptide 2, described on pages 40-41 of the above-captioned patent application, is a smaller polypeptide in which the TRP-LE leader is truncated, a dipeptide Asp-Leu hinge, a Glu cleavage site, and salmon calcitonin analog as a target polypeptide. The salmon calcitonin analog was used because it contained an Asp at position 15 instead of the cleavable native Glu. The sequence of polypeptide 2 is set forth on page 41 of the specification. As stated in the specification on page 41, polypeptide 2 was designed with the goal of obtaining a fusion polypeptide which was cleavable into only 2 fragments, the hook/hinge fragment and calcitonin. Also, the hinge of polypeptide 2 was designed to facilitate cleavage with V-8 protease such that cleavage approached 100% completion.
- 5. Exhibits A and B, attached hereto, demonstrate the experiments in which polypeptides 1 and 2, respectively, were cleaved with V-8 protease. These experiments were performed and the pages submitted as Exhibits A-E are dated prior to March 28, 1986. Exhibits C, D, and E show the results of these cleavage experiments. Exhibit C shows cleavage of polypeptide 1, in which V-8 protease cleaved the fusion protein into multiple fragments. Exhibit D shows cleavage of polypeptide 2, in which V-8 protease cleaved the fusion protein into the two expected fragments, identified as "hook" (containing the leader and hinge regions) and "SC" (containing the salmon calcitonin analog). Exhibit E

shows the purified salmon calcitonin analog collected from the "SC" peak shown in Exhibit D.

6. All statements made herein of our own knowledge are true and all statements made on information and belief are believed to be true; and further these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1011 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

9	29	1/9	12	_
Date	,	7		

9/28/92 Date

9/35/92 Date

9-25-92 Date

9-29-92 Date

9-28-92 Date

9-25-92 Date

8-20-92 Date David C. Rueger

Marc F. Charette

Charles M. Cohen

Poherto Crea

Joines X1

James S. Huston

Peter C Keck

Hermann Oppermann

Richard J. Ridge

Richard J. Ridge

MAR 16 '92 13:02 CBM
ROYECT Calcitonin EXHIBIT A Committe polypapide 1 Continued From Page. 170 130 pm (4133 Ax -24-+1 1:30 ph 9/12/ (C/33-A,15-2/17) 171 13/84 = 7.5% of total = 7/mg = 58.8%

Read and Understood By

Continued on Page

C.H.

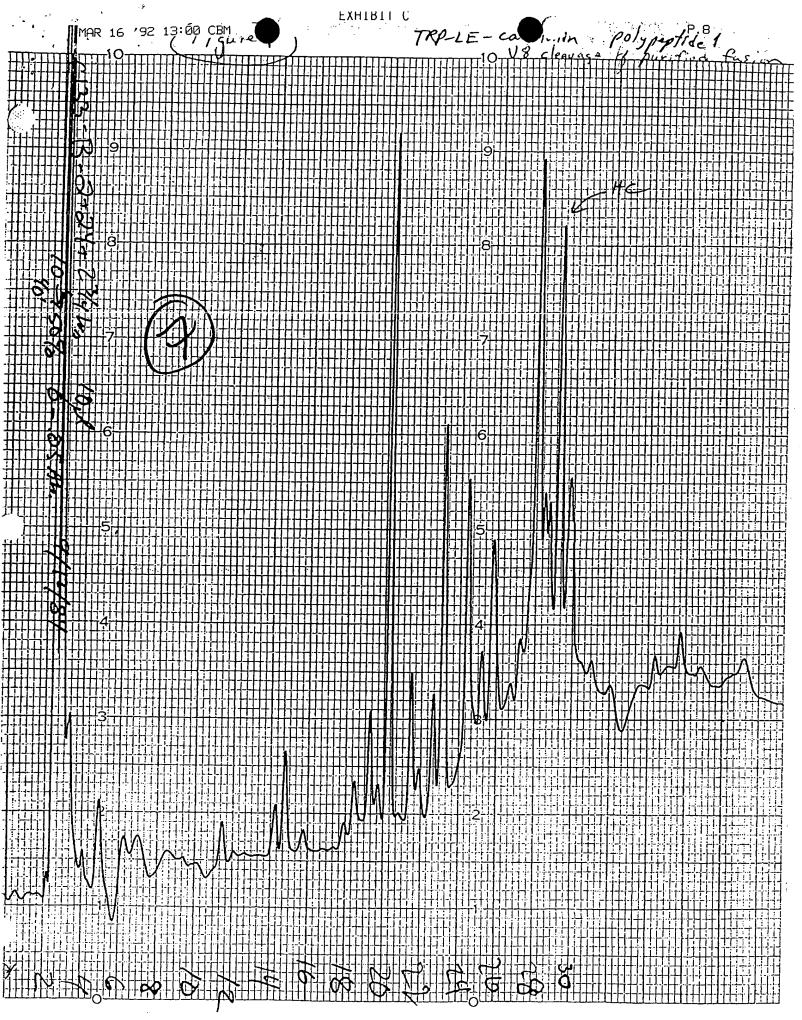
Vita A. Babickas

Continued From Page_____

 -						Т	T	T	T	Τ	1	Π	П	Т	T	Τ	Т	Т	T	T	T	T	T	Т			T	Т				
ч		, , .	ı	r.)	111	+-	۱.۲	+-	╂	+	-	├	-	+-	\vdash	+-	+	 	-	╁,	╁		_	╂	\vdash	-		\vdash		-		
\dashv				P	W	¥	77	er	n	r	a	109	<u> </u>	p^	4-0	Pa	40	4	Py	加	#	50		14	<u> </u>	\vdash	-	-	-	-		
+				17	-	-	 	 	├			<u> </u>	-	}	-	-	+_	-	-		<u> </u>	_	 	 _	_	<u> </u>		-2				
-				K	pl	109		ro			78		0	مر	100	le	12	pw	72	VV	1-4	UX	7	₽_1		1/0	\leq	2/	10	_		
_			Ş	Dù	sli	he.	1/4	pe	Km	1	1	2	$\perp L$	K_	6	Yes	YC	2	المر(إ	1	Hy	Hee	1-6	1/	pl	1 -	8	<u> </u>	(1.	3n	19	ne)
_					<u> </u> '	Ψ_	<u> </u>	<u> </u>	(<u>Y</u>					_		<u> </u>			<u> </u>	Ĺ			<u> </u>	1			<u> </u>				_1
4/ =	7/	85																														
<u>.</u>].																					İ				<u>. </u>							
	A			Voc	54	20		ord	lein	6	esu	ξ	mo	Ine	1	3	Del	2	Dw/	1 1)th	Ac	+1	F. 4	H:	3 ـ ﴿	>					
						+0	12%	pro	1	1	17	1100	U	104	1 /	01	1/	00	sico.			9	5	Б		Cj.	• 7	2 11	c [1		Ţ
\neg						Car	Wa	de	7		45	(+	11	H l			<u> </u>		122									9	ןכנ		
十		\neg				10	The second	1	-	<u>۲</u>	د ر		•	 ' '	7-					-			_								一	\dashv
\dashv	┪	\dashv			-		 							\vdash		 	-														\dashv	十
\dashv	ᅱ	\dashv		-	-	-	-	-		<u> </u>				├	-	-	-				-	-							\dashv	\dashv	-	\dashv
-	3	$ \leftarrow $				n							_		-	-	-				_			-								\dashv
4		4		200	سرو	. 0		ero	Kh	_(رع(2 m	9)	1	100	سر	[2	DW		<i>/</i> /	A	et	E_{i}	ρ	H9							+
	4			士	2	7	M	ero x k	tap	4	Pig	ne	عك	Yuz.	4		1:	12	2	LU	رم		0	12	10						_	_
	\dashv	_			4	1	les	K.		ટ	34	2	\mathcal{L}	1r	+	1/	2	2													1	\rightarrow
<u> </u>											<u>. </u>																					_
			.																													
	<u> </u>)		20	٦,	1 3	10	72Y	· .	7:	1	7	+	in	Du	1 7	Ŋ.,	41	14	4	. +	E,	0	H	×	\neg						\top
1	7			-12	4	20	\exists		~/ ₁		1)	7	,		٠						ر د	, ,		اد.							寸
_	1	一			7,0	×		1 Y	HO			esta	11	J	4	1/2	h,	<u> </u>	يعر	lea	1	٦	-	201				\dashv	一	十	_	十
\dashv	\dashv	\dashv				4	\mathcal{L}	rst Lx lecs	4	9	احـ	۲	_//	7	/	12	MA		\dashv			-		\dashv		-			十		-	\dashv
+	+				-																					\dashv			\dashv	\dashv	+	+
7/2	+	+	-		_	,	7	_	. ,	\dashv			-							_	,			-				-	-		╧┼	-
A_{t}]	\perp		20	7	_4			44			10	$\overline{}$	_	20	W	4	///	ት /	tou	/	5	7	19					_	\dashv	_	\dashv
\perp	1		_		- i) 19	لا	4	ph		Pie	al	<u>.</u>	92	1/4	۱(۲	_4	<u>. 2</u>	22) e	100	٥	@	4.	<u>3</u> \$				_	_	_	_
	\perp				}	4	4	-	70	2	a	14	_	~ 4	26	6														\bot	_	\perp
						1					{	[_							_]	[_ [
Yie	j.H	4	-	,91	2W	. /	17	5	7	(3)	lea	tin		m	. 11		(10	OL	o	10	sch	. 1	42	X						T	$\overline{}$	
丰	Ŧ	>			70				70	$ \tau$		26/	\neg			20	200	1	4					7								T
1	†	-	\dashv	\dashv	f	7	4	- 1	q				3C		90	- 1	1 .1.	- 1	\dashv	+	_	\dashv	\dashv	\dashv	+	+	\dashv	一	十	\dashv	\dashv	+
\dashv	\dagger	\dashv	\dashv	\dashv	:	\dashv	-1	•	44		d		٢	/-	28	\dashv	ell	0.	\dashv	\dashv			\dashv		\dashv	\dashv	-	\dashv	\dashv	\dashv	+	+
+	+	\dashv	\dashv	\dashv	+	\dashv		\dashv	\dashv		, ~				-	$\overline{}$	\dashv	 	-	/			-	\dashv		\dashv	\dashv		\dashv	\dashv	+	+:
+	+	+	\dashv	_	\dashv	-		-	4	-{	"	7	-	56	=	/	91	4	<u>e </u>	 					-+	\dashv		-+	\dashv	+	+	+
+	+	-	+			\dashv	\dashv	\dashv	+	\dashv	#	4	=			4	- 			1					-	\dashv		-+			+	+
- -	\downarrow		\dashv	4	\dashv			_		_		7		\Box	\exists			#	<u> </u>	\dashv			_		_							_L
						1	}									- 1										(Cont	inue	d on	Page	•	

Read and Understood By

C.H.



TRPLE MAR 16 192 13:03 CBM

Polypeptide 2

V8 clearge of marticle fusion EXHIBIT D The identity of the calcitonin analog released from polypeptide 1 after V-8 protease cleavage has been confirmed by sequence analysis to be Phe-Met-Arg-calcitonin. The chromatogram from HPLC analysis of such a digest on a C-18 column developed with a acetonitrile/trifluoroacetic acid (0.1% TFA) gradient is complex since the fusion protein has a molecular weight of about 124,000 daltons (leader approximately 120,000 daltons, hinge calcitonin, approximately 3500 daltons) and contains numerous Glu residues. The calcitonin analog was shown by Western blot analysis to correspond to a specific fraction in the HPLC separation. The fraction was then rechromatographed on a C-8 column yielding pure Phe-Met-Arg-calcitonin.

Isolated Phe-Met-Arg-calcitonin can be cleaved to yield free calcitonin using endo-Arg C protease. The calcitonin can then be isolated by C-18 column chromatography. Its identity has been confirmed by sequence analysis and tryptic peptide mapping.

Example III

A series of fusion polypeptides comprising calcitonin and calcitonin analogs linked to hook polypeptides of various designs were expressed in <u>E. coli</u>. Each fusion protein had designed hook and hinge regions. Each hook contained a glutamic acid residue immediately preceding the N-terminal cysteine residue of calcitonin. The calcitonin was designed to be released from the fusion by cleavage at the Glu residue by V-8 protease.

Polypeptide I had a TRP-LE sequence (188 residues) as a leader, connected at its carboxyl end to a hinge (Phe-Pro-Gly) region, followed by the glutamic acid residue and human calcitonin. It was

hypothesized that this hinge tripeptide would promote cleavage at the Glu residue by providing a flexible residue (proline) and an uncharged "spacer" residue next to the Glu residue. The large leader (27,000 daltons) was used to promote high expression levels. Small leader sequences such as a 17-residue truncated TRP-LE resulted in no expression.

Polypeptide 2 had a modified truncated TRP-LE sequence comprising in sequence a truncated TRP-LE (15 residues) followed by 43 residues derived from the TRP leader sequence (which had been modified by conventional techniques to change the native 3 Met residues to Val, the native 4 Glu residues to Asp, and the single Cys residue to Ser), Asp-Leu as the hinge, the Glu cleavage site, and a salmon calcitonin analog. The salmon calcitonin analog had an Asp at position 15 instead of the native Glu. The internal Glu was discovered to be easily cleaved by V-8 protease. Construction of a calcitonin analog sought to obviate this difficulty. A Gln 15 analog was initially constructed but the fusion expression level was very low. An Asp 15 analog constructed subsequently demonstrated high expression levels.

The structures of polypeptide 1 and 2 are set forth below, the underlined residues indicating desired cleavage sites, the overlined residues indicating the hinge, HC indicating human calcitonin, and SC the salmon calcitonin analog.

Polypeptide 1

MET LYS ALA ILE PHE VAL LEU LYS GLY SER LEU ASP ARG ASP LEU ASP SER ARG ILE GLU LEU GLU MET ARG THR ASP HIS LYS GLU LEU SER GLU HIS LEU MET LEU VAL ASP LEU ALA ARG ASN ASP LEU ALA ARG ILE CYS THR PRO GLY SER ARG TYR VAL ALA ASP LEU THR LYS VAL ASP ARG TYR SER

TYR VAL MET HIS LEU VAL SER ARG VAL VAL GLY GLU LEU
ARG HIS ASP LEU ASP ALA LEU HIS ALA TYR ARG ALA CYS
MET ASN MET GLY THR LEU SER GLY ALA PRO LYS VAL ARG
ALA MET GLN LEU ILE ALA GLU ALA GLU GLY ARG ARG
GLY SER TYR GLY GLY ALA VAL GLY TYR PHE THR ALA HIS
GLY ASP LEU ASP THR CYS ILE VAL ILE ARG SER ALA LEU
VAL GLU ASN GLY ILE ALA THR VAL GLN ALA GLY ALA GLY
VAL VAL LEU ASP SER VAL PRO GLN SER GLU ALA ASP GLU
THR ARG ASN LYS ALA ARG ALA VAL LEU ARG ALA ILE ALA

Polypeptide 2

MET LYS ALA ILE PHE VAL LEU LYS GLY SER LEU ASP ARG ASP LEU ASP SER ARG LEU ASP LEU ASP VAL ARG THR ASP HIS LYS ASP LEU SER ASP HIS LEU VAL LEU VAL ASP LEU ALA ARG ASN ASP LEU ALA ARG ILE VAL THR PRO GLY SER ARG TYR VAL ALA ASP LEU GLU SC

Smaller hook sequences were also designed, but resulted in little or no expression of the fusion polypeptides. Polypeptide I demonstrated good expression levels (15-20%), but because of the presence of multiple Glu and Cys residues in the leader sequence, the post-digestion mixture was complex, containing many fragments, including disulfide linked combinations of fragments. Thus, it was desirable to design a fusion polypeptide that was as small as possible to achieve good expression and could be cleaved into only 2 fragments (the hook/hinge fragment and calcitonin) by V-8 protease. Polypeptide 2 was designed with that goal in mind.